



Single Nucleotide Polymorphism



NEW

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PubMed	Nucleotide	Protein	Genome	Structure	PopSet	Taxonomy
Search	SNP	<input type="text"/>	for			
Limits Preview Index History Clipboard Details						

GENERAL

Publication Detail

NEW

DOCUMENTATION

SEARCH

NEW

Submitter Handle: CGAP-GAI
PMID: [10080189](#)
MEDUID: [99178274](#)
TITLE: Reliable identification of large numbers of candidate SNPs from public EST dat
AUTHOR: [BUETOW,K.H.](#); [EDMONSON, M.N.](#); [CASSIDY, A.B.](#)
JOURNAL: Nat. Genet.
VOLUME: 21
SUPPL: .
ISSUE: .
I_SUPPL: .
PAGES: 323-5
YEAR: 1999
STATUS: 4-published

[View SNP linked to this publication](#)

NEW

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HAPLOTYPE

NEW



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Search SNP <input type="text"/> for <input type="text"/>						
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GENERAL

Submitted SNP(ss) Details

NEW

SNP: Handle|local_snp_id: CGAP-GAI | 52185
NCBI Assay Id(ss#): 10597
Reference SNP Id(rs#): 8598

DOCUMENTATION **STS Information:** Not submitted

Batch Detail:

Submitter Handle: CGAP-GAI
Submitter Batch ID: CANDIDATES.UG.4.20.99
Release Date: Aug 23 1999 5:00PM
Molecular type: cDNA
No. of Chromosomes sampled: 2
Synonym defined:
Organism: Homo sapiens
Submitter Method ID: DEMIGLACE
Citation:

Reliable identification of large numbers of candidate SNPs from data

[View citation details](#)

SEARCH

NEW

SubSNP Detail:

NCBI Assay ID: 10597
Submitter SNP ID: 52185
Synonyms:
LOCUSID: Not available
Submitter STS ID: Not submitted
STS Accession: not available
GenBank Accession: NM 002741 D26181
Comment:

Candidate SNP predicted from aligned EST sequences.
UniGene cluster ID from 4/20/99 build: Hs.2499

HAPLOTYPE **NEW**

Gene Name: PRKCL1
Length: 201
Linkout_URL: <http://lpgws.nci.nih.gov:82/perl/gettrace.p>

Flanking Sequence Information:

5' Flank: TTGTGCCCAC GCTGTCCGGC CGCACCGACG TCAGCAACTT CGACGAGG
AGGCCCCCAC ACTGAGCCCG CCCC GCGACG CGCGGCCCCCT

Observed: C/T

3' Flank: ACAGCCGCGG AGCAGGCAGC CTTCTGGAC TTCGACTTCG TGGCCGGG
cctccccctgc ccctgcccct gccctgccc GAGAGCTCTT

Allele Frequency Information:

POP Batch Id: VALIDATED_01:

Submitter Handle: CGAP-GAI
Submitter Method ID: MASS SPEC
Citation:

High-throughput development and characterization of a genomewide
of gene-based single nucleotide polymorphism markers by chip-based
matrix-assisted laser desorption/ionization time-of-flight mass

Handle|PopulationID: CGAP-GAI|POOLED CEPH
No. of Chromosomes Sampled: 188

Allele: C = 0.562 / T = 0.438

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DOCUMENTATION: [FAQ](#) | [Overview](#) | [How To Submit](#) | [RefSNP Summary Info](#) | [Database Schema](#)

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PubMed	Nucleotide	Protein	Genome	Structure	PopSet	Taxonomy
Search SNP <input type="text"/> for <input type="text"/>						
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GENERAL

Submitted SNP(ss) Details

NEW

SNP: Handle|local_snp_id: CGAP-GAI | 56405
NCBI Assay Id(ss#): 13018
Reference SNP Id(rs#): 10846

DOCUMENTATION

STS Information: Not submitted

Batch Detail:

Submitter Handle: CGAP-GAI
Submitter Batch ID: CANDIDATES.UG.4.20.99
Release Date: Aug 23 1999 5:00PM
Molecular type: cDNA
No. of Chromosomes sampled: 2
Synonym defined:
Organism: Homo sapiens
Submitter Method ID: DEMIGLACE
Citation:

Reliable identification of large numbers of candidate SNPs from data

[View citation details](#)

SEARCH

NEW

SubSNP Detail:

NCBI Assay ID: 13018
Submitter SNP ID: 56405
Synonyms:
LOCUSID: Not available
Submitter STS ID: Not submitted
STS Accession: not available
GenBank Accession: NM 002741 D26181
Comment:

Candidate SNP predicted from aligned EST sequences.
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Observed: G/A

3' Flank: TCAGCAACTT CGACGAGGAG TTCACCGGGG AGGCCCCCAG ACTGAGCC
CGCGGGCCCT TACAGCCGCG GAGCAGGCAG CCTTCCTGGA

Allele Frequency Information:

POP Batch Id: VALIDATED_01:

Submitter Handle: CGAP-GAI
Submitter Method ID: MASS SPEC
Citation:

High-throughput development and characterization of a genomewide
of gene-based single nucleotide polymorphism markers by chip-based
matrix-assisted laser desorption/ionization time-of-flight mass

Handle|PopulationID: CGAP-GAI|POOLED CEPH
No. of Chromosomes Sampled: 188

Allele: A = 0.176 / G = 0.824

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